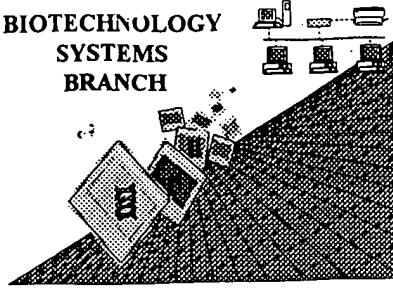


0590  
1005

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/894,912  
Source: OIPK  
Date Processed by STIC: 7/18/2001

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**  
**<http://www.uspto.gov/web/offices/pac/checker>**

OIPE

**RAW SEQUENCE LISTING** DATE: 07/18/2001  
**PATENT APPLICATION:** US/09/894,912 TIME: 10:30:01

Input Set : A:\37260Aseq.txt  
Output Set: N:\CRF3\07182001\I894912.raw

5 <110> APPLICANT: Tang et al.  
8 <120> TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO NOVEL STEM CELL GROWTH  
FACTOR-LIKE

9 POLYPEPTIDES AND POLYNUCLEOTIDES  
12 <130> FILE REFERENCE: 28110/37260A  
~~14~~ > **14 <140> CURRENT APPLICATION NUMBER: US/09/894,912**

15 <141> CURRENT FILING DATE: 2001-06-28  
17 <150> PRIOR APPLICATION NUMBER: To be assigned  
18 <151> PRIOR FILING DATE: 2001-04-05  
20 <150> PRIOR APPLICATION NUMBER: 60/266,614  
21 <151> PRIOR FILING DATE: 2001-02-05  
23 <150> PRIOR APPLICATION NUMBER: 60/215,733  
24 <151> PRIOR FILING DATE: 2000-06-28  
26 <150> PRIOR APPLICATION NUMBER: 09/757,562  
27 <151> PRIOR FILING DATE: 2001-01-09  
29 <150> PRIOR APPLICATION NUMBER: 09/543,774  
30 <151> PRIOR FILING DATE: 2000-04-05  
33 <160> NUMBER OF SEQ ID NOS: 48  
35 <170> SOFTWARE: PatentIn version 3.0  
39 <210> SEQ ID NO: 1  
40 <211> LENGTH: 301  
41 <212> TYPE: DNA  
42 <213> ORGANISM: Homo sapiens  
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45	gcacgagacg aggaaaaaaaaa ggaagggaga ggaaaaagaaa aaaacctaatt aaaggagaaaa	60
47	gtaaaagaagc aataccgtac agcaaaagtc tggatccag caaagaaaatc ccagagcaac	120
49	gagaaaaacaa acagcagcag aagaagcgaa aagtccaaga taaacagaaaa tcggtatcag	180
51	tcagcactgt acactagagg gttccatgag attattgttag actcatgtatctg ctgttatctc	240
53	aaccagatgc ccaggacagg tgctctagcc attaggacca caaatggaca tgtcagttat	300
55	t	301
58	<210> SEQ ID NO: 2	
59	<211> LENGTH: 392	
60	<212> TYPE: DNA	
61	<213> ORGANISM: Homo sapiens	
63	<400> SEQUENCE: 2	
64	tggaaactcga tatccagata taaataagcg tacaaaatgc aaagctgact gtgataccgt	60
66	tttcaacaaa gatttctgca caaaatgtaa aagtggattt tacttacacc ttggaaagtgc	120
68	ccttgacaat tgcccagaag gtttggaaagc caacaaccat actatggagt gtgtcagtat	180
70	tgtgcactgt gaggtcagtg aatggaaatcc ttggagtcca tgcacgaaga agggaaaaaac	240
72	atgtggcttc aaaagaggaa ctgaaaacacg gttccgagaa ataatacagc atccttcagc	300
74	aaagggttaac ctatgtcccc caacaaatga gacaagaaag tgtacagtgc aaaggaagaa	360
76	gtgtcagaag ggagaacgag gaaaataagg ag	392
79	<210> SEQ ID NO: 3	
80	<211> LENGTH: 475	
81	<212> TYPE: DNA	
82	<213> ORGANISM: Homo sapiens	
84	<220> FEATURE:	
85	<221> NAME/KEY: misc_feature	

*Does Not Comply  
Corrected Diskette Needed*

*M 6/7*

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/894,912

DATE: 07/18/2001  
TIME: 10:30:01

Input Set : A:\37260Aseq.txt  
Output Set: N:\CRF3\07182001\I894912.raw

87 <222> LOCATION: (1)..(475)  
 88 <223> OTHER INFORMATION: n = A, T, G, or C  
 91 <400> SEQUENCE: 3

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W--> 94 ccacgcgtnc gcccacgcgt ccggagagga aaagaaaaaa acctaattta ggagaaaagta	120
96 aagaagcaat acctgacagc ggaagtctgg aatggagcaa agaaatccca gagcaacgag	180
98 aaaacaaaca gcagcagaag aagcgaaaag tccaagataa acagaaatcg gtatcgtca	240
100 gcactgtaca cttagagggtt ccatgagatt atttagact catgatgtc ctatctaac	300
102 cagatgccca ggacaggtgc tctagccatt aggaccacaa atggacatgt cagttattgc	360
104 tctgtctaaa caacattccc agtagttgtc atattcttca tacaagcata gttaacaaca	420
W--> 106 aagagccaaa agatcaaaga agggatactt tcagatgggt gtcttgtgtc ttctn	475
109 <210> SEQ ID NO: 4	
110 <211> LENGTH: 473	
111 <212> TYPE: DNA	
112 <213> ORGANISM: Homo sapiens	
114 <220> FEATURE:	
115 <221> NAME/KEY: misc_feature	
117 <222> LOCATION: (1)..(473)	
118 <223> OTHER INFORMATION: n = A, T, G, or C	
121 <400> SEQUENCE: 4	
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124 aaaaaaggaa gggagaggaa aagaaaaaaaaa cctaataaaag gagaaagtaa agaagcaata	120
126 cctgacagca aaagtctgga atccagcaga gaaatcccg agcaacgaga aaacaaacag	180
128 cagcagaaga agcgaaaagt ccaagataaa cagaaatcg tatcagtctac cactgtacac	240
130 tagagggttc catgagatta ttgttagactc atgatgtc tatctcaacc agatgccag	300
132 gacaggtgtc cttagccatta ggaccacaaa tggacatgtc agttattgtc ctgtctaaac	360
134 aacattccca gtagttgcta tattcttcat acaagcatag ttaacaacaa agagccaaaa	420
136 gatcaaagaa gggatacttt cagatgggtc ttcttgtgc ttctctgtcat ttt	473
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140 <211> LENGTH: 462	
141 <212> TYPE: DNA	
142 <213> ORGANISM: Homo sapiens	
145 <220> FEATURE:	
146 <221> NAME/KEY: misc_feature	
148 <222> LOCATION: (1)..(462)	
149 <223> OTHER INFORMATION: n = A, T, G, or C	
152 <400> SEQUENCE: 5	
W--> 153 tgggagannn nttgaaact gagatcgatc canacncnac nangaataaa aggaagggag	60
155 agggaaagaa aaaaacctaa taaaggagaa agttaagaat caatttctga cagcaaaagt	120
157 ctggaatcca tcaaagaaat cccatatcaa cgagaaaaca gacagcagca caaaaagcga	180
159 aaagtccaaat ataaacagaa atcggtatca gtcagactg tacactagag gttccatga	240
161 gattattgtt gactcatgat gctgttatct caaccagatg cccaggacag gtgtctatc	300
163 cattacgacc acaaatggac atgtcgttta ttgtctgtc taaacaacat tcccgatgt	360
165 tgctatattt ttcataacaag catagttaa aacaaagagc caaaaagatca aagaagggat	420
167 actttcagat gttgtcttg tgtgttctc tgcatatata aa	462
170 <210> SEQ ID NO: 6	
171 <211> LENGTH: 384	
172 <212> TYPE: DNA	
173 <213> ORGANISM: Homo sapiens	

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/894,912**

**DATE: 07/18/2001**  
**TIME: 10:30:02**

**Input Set : A:\37260Aseq.txt**  
**Output Set: N:\CRF3\07182001\I894912.raw**

176 <400> SEQUENCE: 6  
 177 aataatgtgt aacaaatgca aagctgactg tgataacctgt ttcaacaaaa atttctgcac 60  
 179 aaaatgtaaa agtgatttt acttacacct tggaaagtgc cttgacaatt gcccagaagg 120  
 181 gttgaagacc aacaaccata ctatggagtg tgtcagtatt gtgcactgtg aggtcagtga 180  
 183 atggaatcct tggagtccat gcacgaagaa gggaaaaaca tgtggcttca aaagagggac 240  
 185 tggaaacacgg gtccgagaaa taatacagca tccttcagca aagggttaacc tatgtcccc 300  
 187 aacaaatgag acaagaaaagt gtacagtgc aaggaagaag tgtcagaagg gagaacgagg 360  
 189 aaaaaaagga agggagagga aaag 384  
 192 <210> SEQ ID NO: 7  
 193 <211> LENGTH: 390  
 194 <212> TYPE: DNA  
 195 <213> ORGANISM: Homo sapiens  
 197 <220> FEATURE:  
 198 <221> NAME/KEY: misc\_feature  
 200 <222> LOCATION: (1)..(390)  
 201 <223> OTHER INFORMATION: n = A, T, G, or C  
 204 <400> SEQUENCE: 7  
 205 cgttgctctg ggatttcttt gctggattcc agactttgc tgtcaggat tgcttcttta 60  
 207 ctttctcctt tattaggttt ttttctttc ctctcccttc ctttttttcc tcgttctccc 120  
 209 ttctgacact ttttcttgc cactgtacac ttttctgtct catttgttgg gggacatagg 180  
 211 ttaccctttt ctgaaggatg ctgtattatt tctcgaccc gtgtttcagt ccctctttg 240  
 213 aagccacatg ttttccctt ctctgtgc cat ggactccaag gattccattc actgaccta 300  
 215 cagtgcacaa tactgacaca ctccatagta tgggttgttgg cttccaaccc ttctggcaa 360  
 217 **ttgtcaaggc acttccaag gtgtaa** 390  
 220 <210> SEQ ID NO: 8  
 221 <211> LENGTH: 1345  
 222 <212> TYPE: DNA  
 223 <213> ORGANISM: Homo sapiens  
 225 <220> FEATURE:  
 226 <221> NAME/KEY: misc\_feature  
 228 <222> LOCATION: (321)..(1235)  
 229 <223> OTHER INFORMATION: similar to gi4519541 in the genpept database release 114,

Run wit

230 h FASTXY3.3t00, default parameter  
 233 <400> SEQUENCE: 8  
 234 gccccggccc cggccgctcc tggAACCCG gttcgccgc atgccagcca ccccagcgaa 60  
 236 gcccggcag ttcatgtctt ggataatttg aaagtacaat agttggttc cctgtccacc 120  
 238 cggcccaact cgcttgccat cacagcacgc ctatcgatg tgagaggaga agtcccgctg 180  
 240 ctcgggact gtctatatac gcctaacc tacatatatt taaaaacat taaatataat 240  
 242 taacaatcaa aagaaagagg agaaaggaa ggaagcatta ctgggttact atgcacttgc 300  
 244 gactgatttc ttggctttt atcatttga actttatgaa atacatcgcc agccaaaacg 360  
 246 cttccgggg aaggcgccag cgaagaatgc atcctaactgt tagtcaaggc tgccaaggag 420  
 248 gctgtcaac atgctcagat tacaatggat gtttgtcatg taagcccaga ctatTTTGT 480  
 250 ctctggaaag aattggcatg aagcagattg gagatgtct catcttcatg tccaagtgg 540  
 252 tattatggaa ctcgatatac agatataaat aatgtgtaca aaatgcaaag ctgactgtga 600  
 254 tacctgtttc aacaaaaatt tctgcacaaa atgtaaaagt ggattttact tacacccttgg 660  
 256 aaagtgcctt gacaattgcc cagaagggtt ggaagccac aaccatacta tggagtgtgt 720  
 258 cagtattgtg cactgtgagg tcagtgaatg gaatccttgg agtccatgca cgaagaagg 780  
 260 aaaaacatgt ggcttcaaaa gagggactga aacacgggtc cgagaaataa tacagcatcc 840  
 262 ttcagcaaag ggtaaacctat gtccccaac aaatgagaca agaaagtgt a cagtgcacaa 900

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/894,912

DATE: 07/18/2001

TIME: 10:30:02

Input Set : A:\37260Aseq.txt

Output Set: N:\CRF3\07182001\I894912.raw

264	gaagaagtgt	cagaaggggag	aacgaggaaaa	aaaaggaagg	gagaggaaaa	gaaaaaaaaacc	960
266	taataaaagga	gaaagtaaaag	aagcaatacc	tgacagcaaa	agtctggaat	ccagcaaaga	1020
268	aatcccagag	caacgagaaa	acaaacagca	gcagaagaag	cgaaaagtcc	aagataaaaca	1080
270	gaaatcggtt	tcagtcagca	ctgtacacta	gagggttcca	tgagattatt	gtagactcat	1140
272	gatgctgcta	tctcaaccag	atgcccagga	caggtctct	agccattagg	accacaaatg	1200
274	gacatgtcag	ttattgctct	gtctaaacaa	cattcccagt	agttgctata	ttcttcatac	1260
276	aagcatagg	aacaacaaag	agccaaaaga	tcaaagaagg	gatacttca	gatgggtgtc	1320
278	ttgtgtgctt	ctctgcattt	ttaaa				1345
281	<210>	SEQ ID NO:	9				
282	<211>	LENGTH:	1343				
283	<212>	TYPE:	DNA				
284	<213>	ORGANISM:	Homo sapiens				
286	<220>	FEATURE:					
287	<221>	NAME/KEY:	CDS				
289	<222>	LOCATION:	(291)..(1109)				
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295	gccgcgcag	ttcagtgttt	ggataatttg	aaagtacaat	agttgggttc	cctgtccacc	120
297	cgcggccactt	cgcttgccat	cacagcacgc	ctatcgatg	tgagaggaga	agtcccgtg	180
299	ctcgggcact	gtcttatatac	gcctaacacc	tacatatatt	ttaaaaacat	taaatataat	240
301	taacaatcaa	aagaaagagg	agaaaggaag	ggaagcatta	ctgggttact	atg cac	296
302					Met His		
303					1		
305	ttg cga ctg	att tct tgg	ctt ttt atc	att ttg aac	ttt atg gaa	tac	344
306	Leu Arg	Leu Ile Ser	Trp Leu Phe	Ile Ile Leu Asn	Phe Met	Glu Tyr	
307	5	10	15				
309	atc ggc agc	caa aac gcc	tcc cgg gga	agg cgc cag	cga aga atg	cat	392
310	Ile Gly	Ser Gln Asn	Ala Ser Arg	Gly Arg Arg	Gln Arg Arg	Met His	
311	20	25	30				
313	cct aac gtt	agt caa ggc	tgc caa gga	ggc tgt gca	aca tgc tca	gat	440
314	Pro Asn Val	Ser Gln Gly	Cys Gln Gly	Gly Cys Ala	Thr Cys Ser	Asp	
315	35	40	45				
319	tac aat gga	tgt ttg tca	tgt aag ccc	aga cta ttt	ttt gct ctg	gaa	488
320	Tyr Asn Gly	Cys Leu Ser	Cys Lys Pro	Arg Leu Phe	Phe Ala	Leu Glu	
321	55	60	65				
323	aga att ggc	atg aag cag	att gga gta	tgt ctc tct	tca tgt cca	agt	536
324	Arg Ile Gly	Met Lys Gln	Ile Gly Val	Cys Leu Ser	Ser Cys Pro	Ser	
325	70	75	80				
327	gga tat tat	gga act cga	tat cca gat	ata aat aag	tgt aca aaa	tgc	584
328	Gly Tyr	Tyr Gly Thr	Arg Tyr Pro	Asp Ile Asn	Lys Cys Thr	Lys Cys	
329	85	90	95				
331	aaa gct gac	tgt gat acc	tgt ttc aac	aaa aat ttc	tgc aca aaa	tgt	632
332	Lys Ala Asp	Cys Asp Thr	Cys Phe Asn	Lys Asn Phe	Cys Thr	Lys Cys	
333	100	105	110				
335	aaa agt gga	ttt tac tta	cac ctt gga	aag tgc ctt	gac aat tgc	cca	680
336	Lys Ser Gly	Phe Tyr Leu	His Leu Gly	Lys Cys Leu	Asp Asn	Cys Pro	
337	115	120	125				
339	gaa ggg ttg	gaa gcc aac	aac cat act	atg gag	tgt gtc	agt att gtg	728
340	Glu Gly Leu	Glu Ala Asn	His Thr Met	Glu Cys Val	Ser Ile	Val	

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/894,912**

**DATE: 07/18/2001**  
**TIME: 10:30:02**

**Input Set : A:\37260Aseq.txt**  
**Output Set: N:\CRF3\07182001\I894912.raw**

341	135	140	145	
343	cac tgt gag gtc agt gaa tgg aat cct tgg agt cca tgc acg aag aag			776
344	His Cys Glu Val Ser Glu Trp Asn Pro Trp Ser Pro Cys Thr Lys Lys			
345	150	155	160	
347	gga aaa aca tgt ggc ttc aaa aga ggg act gaa aca cgg gtc cga gaa			824
348	Gly Lys Thr Cys Gly Phe Lys Arg Gly Thr Glu Thr Arg Val Arg Glu			
349	165	170	175	
351	ata ata cag cat cct tca gca aag ggt aac cta tgt ccc cca aca aat			872
352	Ile Ile Gln His Pro Ser Ala Lys Gly Asn Leu Cys Pro Pro Thr Asn			
353	180	185	190	
355	gag aca aga aag tgt aca gtg caa agg aag aag tgt cag aag gga gaa			920
356	Glu Thr Arg Lys Cys Thr Val Gln Arg Lys Lys Cys Gln Lys Gly Glu			
357	195	200	205	210
359	cga gga aaa aaa gga agg gag agg aaa aga aaa aaa cct aat aaa gga			968
360	Arg Gly Lys Lys Gly Arg Glu Arg Lys Arg Lys Pro Asn Lys Gly			
361	215	220	225	
363	gaa agt aaa gaa gca ata cct gac agc aaa agt ctg gaa tcc agc aaa			1016
364	Glu Ser Lys Glu Ala Ile Pro Asp Ser Lys Ser Leu Glu Ser Ser Lys			
365	230	235	240	
367	gaa atc cca gag caa cga gaa aac cag cag cag aag aag cga aaa			1064
368	Glu Ile Pro Glu Gln Arg Glu Asn Lys Gln Gln Lys Lys Arg Lys			
369	245	250	255	
371	gtc caa gat aaa cag aaa tcg gta tca gtc agc act gta cac tag			1109
372	Val Gln Asp Lys Gln Lys Ser Val Ser Val Ser Thr Val His			
373	260	265	270	
375	agggttccat gagattattg tagactcatg atgctgctat ctcaaccaga tgcccaggac			1169
377	aggtgctcta gccatttagga ccacaaatgg acatgtcagt tattgctctg tctaaacaac			1229
379	attcccgata gttgctatat tcttcataca agcatagtt acaacaaaga gccaaaagat			1289
381	caaagaaggg atacttcag atgggtgtct tgtgtgcttc tctgcatttt taaa			1343
384	<210> SEQ ID NO: 10			
385	<211> LENGTH: 272			
386	<212> TYPE: PRT			
387	<213> ORGANISM: Homo sapiens			
390	<400> SEQUENCE: 10			
392	Met His Leu Arg Leu Ile Ser Trp Leu Phe Ile Ile Leu Asn Phe Met			
393	1	5	10	15
396	Glu Tyr Ile Gly Ser Gln Asn Ala Ser Arg Gly Arg Arg Gln Arg Arg			
397	20	25	30	
400	Met His Pro Asn Val Ser Gln Gly Cys Gln Gly Gly Cys Ala Thr Cys			
401	35	40	45	
404	Ser Asp Tyr Asn Gly Cys Leu Ser Cys Lys Pro Arg Leu Phe Phe Ala			
405	50	55	60	
408	Leu Glu Arg Ile Gly Met Lys Gln Ile Gly Val Cys Leu Ser Ser Cys			
409	65	70	75	80
412	Pro Ser Gly Tyr Tyr Gly Thr Arg Tyr Pro Asp Ile Asn Lys Cys Thr			
413	85	90	95	
416	Lys Cys Lys Ala Asp Cys Asp Thr Cys Phe Asn Lys Asn Phe Cys Thr			
417	100	105	110	
420	Lys Cys Lys Ser Gly Phe Tyr Leu His Leu Gly Lys Cys Leu Asp Asn			

09/894,912 6

<210> 31  
<211> 2384  
<212> DNA  
<213> Mus musculus

<220>  
<221> CDS  
<222> (511)..(1347)

<400> 31

See next page for erroneous portion of seq. 31

09/894,912 7

agggaaaaac tattttatta cctgcctatg ggtccacctg gaactaaagg gatactactt 2307

tcttaacaagg tgtatctagt aggagagaaa gccaccacaa taaatatatt tgtaatagn 2367

aaaaaaaaa aaaaaaaaaa 2384

see item 9  
on Error  
summary  
sheet



Use of n and/or Xaa has been detected in the Sequence Listing.  
Review the Sequence Listing to insure a corresponding  
explanation is presented in the <220> to <223> fields of  
each sequence using n or Xaa.

**VERIFICATION SUMMARY**  
PATENT APPLICATION: US/09/894,912

DATE: 07/18/2001  
TIME: 10:30:03

Input Set : A:\37260Aseq.txt  
Output Set: N:\CRF3\07182001\I894912.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number  
L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:217 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:1201 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:31  
L:1201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31